

LDIR

Ref #	Hits	Search Query	DBs	Default Operator	Plurals	Time Stamp
L1	43	mutation and hamming and primer	US-PGPUB; USPAT; EPO; JPO; DERWENT	ADJ	ON	2007/05/15 09:37
L2	43	mutation and hamming and primer	US-PGPUB; USPAT; DERWENT	ADJ	ON	2007/05/15 11:39
L3	140	(mismatch near3 tolerance) and primer	US-PGPUB; USPAT; DERWENT	ADJ	ON	2007/05/15 11:34
L4	0	(mismatch near3 tolerance) and primer and occurrence	US-PGPUB; USPAT; DERWENT	ADJ	ON	2007/05/15 11:35
L5	16	(mismatch near3 tolerance) and primer and subsequence	US-PGPUB; USPAT; DERWENT	ADJ	ON	2007/05/15 11:37
L6	14	(mismatch near3 tolerance) and hamming	US-PGPUB; USPAT; DERWENT	ADJ	ON	2007/05/15 11:38
L7	1	(mismatch near3 tolerance) and hamming and DNA	US-PGPUB; USPAT; DERWENT	ADJ	ON	2007/05/15 11:38
L8	43	mutation and hamming and primer and DNA	US-PGPUB; USPAT; DERWENT	ADJ	ON	2007/05/15 11:41
L9	13	mutation and hamming and (primer near3 (selection or method)) and DNA	US-PGPUB; USPAT; DERWENT	ADJ	ON	2007/05/15 12:37
L10	0	k-tuple and hamming and (primer near3 (selection or method)) and DNA	US-PGPUB; USPAT; DERWENT	ADJ	ON	2007/05/15 12:37
L11	45	k-tuple and hamming	US-PGPUB; USPAT; DERWENT	ADJ	ON	2007/05/15 12:37
L12	3	k-tuple and hamming and primer	US-PGPUB; USPAT; DERWENT	ADJ	ON	2007/05/15 12:38
L13	0	k-tuple and hamming and DNA	US-PGPUB;	ADJ	ON	2007/05/15 12:38

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			USPAT; DERWENT			
L14	38	k-tuple and hamming and sequence	US- PGPUB; USPAT; DERWENT	ADJ	ON	2007/05/15 12:38

STN Search 5/15/07

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(FILE 'HOME' ENTERED AT 08:16:59 ON 15 MAY 2007)  
SET PLURALS ON  
SET ABB ON

FILE 'CAPLUS, MEDLINE, SCISEARCH' ENTERED AT 08:17:26 ON 15 MAY  
2007

L1 13 S SUBSEQUENCE AND HAMMING  
L2 126638 S (PROBE OR PRIMER) AND (MUTAT? OR POLYMORPH?)  
L3 81056 S (PROBE OR PRIMER) (L) (MUTAT? OR POLYMORPH?)  
L4 47516 S (PRIMER) (L) (MUTAT? OR POLYMORPH?)  
L5 2090 S HAMMING  
L6 80 S OCCURENCE (3A) FREQUENCY  
L7 0 S L4 AND L5 AND L6  
L8 1 S PRIMER AND L6  
L9 980 S PRIMER (3A) SELECTION  
L10 1 S L5 AND L9  
L11 1861 S PRIMER (S) SELECTION  
L12 1 S L5 AND L11  
L13 6 S HAMMING AND PRIMER

FILE 'CAPLUS, MEDLINE, BIOSIS, EMBASE, SCISEARCH' ENTERED AT  
08:49:47 ON  
15 MAY 2007

L14 7 S L13  
L15 7 DUP REM L14 (0 DUPLICATES REMOVED)  
L16 6 S "CLOSEST STRING PROBLEM"  
L17 100 S SEQUENCE AND SUBSEQUENCE AND PRIMER  
L18 15 S HAMMING AND SUBSEQUENCE  
L19 8 DUP REM L18 (7 DUPLICATES REMOVED)  
L20 89 S HAMMING AND DNA  
L21 57 DUP REM L20 (32 DUPLICATES REMOVED)  
L22 5 S L21 AND MUTAT?  
L23 15 S SUBSEQUENCE AND HAMMING  
L24 8 DUP REM L23 (7 DUPLICATES REMOVED)  
L25 649 S MUTATION AND HAM? AND PRIMER  
L26 1 S MUTATION AND HAMMING AND PRIMER  
L27 9 S MORISHITA S/AU AND PRIMER  
L28 168 S YAMADA T/AU AND PRIMER  
L29 261 S MISMATCH (5A) TOLERANCE  
L30 190 S MISMATCH (3A) TOLERANCE  
L31 156 S OCCURENCE (3A) FREQUENCY  
L32 0 S L30 AND L31

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L33	0 S L30 AND HAMMING
L34	0 S L31 AND HAMMING
L35	19 S L30 AND PRIMER
L36	10 DUP REM L35 (9 DUPLICATES REMOVED)